10/537512

SEQUENCE LISTING JC17 Rec'd PCT/PTO 03 JUN 2005

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Val Ser Phe Ala Leu Pro Ser Ala Glu Lys Asp Met Cys Leu Ser Ser 130 135 140

Ser Lys Lys Gly Met Leu Gly Met Ile Val Tyr Leu Gly Met Met Ala 145 150 155 160

Gly Ala Phe Ile Leu Gly Gly Leu Ala Asp Lys Leu Gly Arg Lys Arg 165 170 175

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Asp Val Thr Ser Thr Asp Thr Tyr Phe Lys Asn Cys Thr Ile Glu Ser 485 490 495

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				ata Ile						1536
				tcc Ser						1584
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				aaa Lys						1680
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Gly Leu Ser Val Trp Phe Pro Asp Val Ile Lys Pro Leu Gln Ser Asp 450 455 460

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agc ggc atg gca aga gtg ggt gct ctc atc act ccg ttc atc gcc cag Ser Gly Met Ala Arg Val Gly Ala Leu Ile Thr Pro Phe Ile Ala Gln 465 470 475 480	1440
gtg atg ctg gaa tcc tct gtg tac ctg act ctg gca gtt tac agt ggc Val Met Leu Glu Ser Ser Val Tyr Leu Thr Leu Ala Val Tyr Ser Gly 485 490 495	1488
tgc tgc ctc ctg gct gcc ctg gcc tcc tgc ttt ttg ccc att gag acc Cys Cys Leu Leu Ala Ala Leu Ala Ser Cys Phe Leu Pro Ile Glu Thr 500 505 510	1536
aaa ggc cga gga ctg cag gag tcc agc cac cgg gag tgg ggc cag gag Lys Gly Arg Gly Leu Gln Glu Ser Ser His Arg Glu Trp Gly Gln Glu 515 520 525	1584
atg gtc ggc cga gga atg cac ggt gca ggt gtt acc agg tcg aac tct Met Val Gly Arg Gly Met His Gly Ala Gly Val Thr Arg Ser Asn Ser 530 535 540	1632
ggc tct cag gaa tag Gly Ser Gln Glu 545	1647
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Glu His Glu Val Gln Ile Glu Gly Val His Val Gly Leu Glu Ala Val 35 40 45	

Asp Asp Thr Phe Met Val Glu Asp Ala Val Glu Ala Ile Gly Phe Gly 65 70 75 80

Lys Phe Gln Trp Lys Leu Ser Val Leu Thr Gly Leu Ala Trp Met Ala 85 90 95

Asp Ala Met Glu Met Met Ile Leu Ser Ile Leu Ala Pro Gln Leu His $100 \hspace{1.5cm} 105 \hspace{1.5cm} 110$

Cys Glu Trp Arg Leu Pro Ser Trp Gln Val Ala Leu Leu Thr Ser Val 115 120 125

Val Phe Val Gly Met Met Ser Ser Ser Thr Leu Trp Gly Asn Ile Ser 130 135 140

Asp Gln Tyr Gly Arg Lys Thr Gly Leu Lys Ile Ser Val Leu Trp Thr 145 150 155 160

Leu Tyr Tyr Gly Ile Leu Ser Ala Phe Ala Pro Val Tyr Ser Trp Ile 165 170 175

Leu Val Leu Arg Gly Leu Val Gly Phe Gly Ile Gly Gly Val Pro Gln
180 185 190

Ser Val Thr Leu Tyr Ala Glu Phe Leu Pro Met Lys Ala Arg Ala Lys 195 200 205

Cys Ile Leu Leu Ile Glu Val Phe Trp Ala Ile Gly Thr Val Phe Glu 210 215 220

Val Val Leu Ala Val Phe Val Met Pro Ser Leu Gly Trp Arg Trp Leu 225 230 235 240

Leu Ile Leu Ser Ala Val Pro Leu Leu Phe Ala Val Leu Cys Phe 245 250 255

Trp Leu Pro Glu Ser Ala Arg Tyr Asp Val Leu Ser Gly Asn Gln Glu 260 265 270

Lys Ala Ile Ala Thr Leu Lys Arg Ile Ala Thr Glu Asn Gly Ala Pro 275 280 285

Met	Pro	Leu	Gly	Lys	Leu	Ile	Ile	Ser	Arg	Gln	Glu	Asp	Arg	Gly	Lys
	290					295					300				

Met Arg	Asp	Leu	Phe	Thr	Pro	His	Phe	Arg	Trp	Thr	Thr	Leu	Leu	Leu
305				310					315					320

Leu Thr Thr Glu Leu Phe Gln Ala Gly Asp Val Cys Gly Ile Ser Ser
$$340$$
 345 350

Ser Glu Glu Asp Tyr Met Asp Leu Leu Trp Thr Thr Leu Ser Glu Phe
$$370$$
 380

Lys Thr Met Ala Leu Cys Phe Val Ile Phe Ser Phe Cys Ser Leu Leu
$$405$$
 410 415

Leu Phe Ile Cys Val Gly Arg Asn Val Leu Thr Leu Leu Phe Ile 420 425 430

Ala Arg Ala Phe Ile Ser Gly Gly Phe Gln Ala Ala Tyr Val Tyr Thr 435 440 445

Pro Glu Val Tyr Pro Thr Ala Thr Arg Ala Leu Gly Leu Gly Thr Cys 450 455 460

Ser Gly Met Ala Arg Val Gly Ala Leu Ile Thr Pro Phe Ile Ala Gln 465 470 475 480

Val Met Leu Glu Ser Ser Val Tyr Leu Thr Leu Ala Val Tyr Ser Gly 485 490 495

Cys Cys Leu Leu Ala Ala Leu Ala Ser Cys Phe Leu Pro Ile Glu Thr 500 505 510

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ttctaaaggg tcttcttgat ccctccaatt cattgagcaa agggctgaaa aagaagcaga	360
gagtaaggta gagccagtga ctcgccccca agccccatc atg gaa gaa ggc ttt Met Glu Glu Gly Phe 1 5	414
cga gac cga gca gcg ttc atc cgt ggg gcc aaa gac att gcc aag gaa Arg Asp Arg Ala Ala Phe Ile Arg Gly Ala Lys Asp Ile Ala Lys Glu 10 15 20	462
gtt aag aag cac gcg gcc aag aag gtg gtg aag ggt ctc gac aga gtc Val Lys Lys His Ala Ala Lys Lys Val Val Lys Gly Leu Asp Arg Val 25 30 35	510
cag gat gaa tat too oga agg too tac too ogo ttt gag gag gag gag Gln Asp Glu Tyr Ser Arg Arg Ser Tyr Ser Arg Phe Glu Glu Glu 40 45 50	558
gat gat gat gac ttc cct gcc cct gct gac ggc tat tac cgc gga gaa Asp Asp Asp Asp Phe Pro Ala Pro Ala Asp Gly Tyr Tyr Arg Gly Glu 55 60 65	606
ggg gcc cag gat gag gaa ggt ggc gct tcc agt gat gcc act gag	654

Gly 70	Ala	Gln	Asp	Glu	Glu 75	Glu	Gly	Gly	Ala	Ser 80	Ser	Asp	Ala	Thr	Glu 85	
					gat Asp											702
					Gly ggg											750
					aga Arg											798
					gcg Ala											846
					atc Ile 155											894
					gtg Val											942
					ggc Gly											990
					aac Asn											1038
					gcc Ala											1086
					ctg Leu 235											1134
					ttc Phe			Gly								1182
					gtt Val											1230
					ttt Phe											1278
agc	tgg	ctc	tgt	atg	ttc	tgg	atg	att	ggt	ggc	gtg	tat	gca	gct	gca	1326

Ser	Trp 295	Leu	Cys	Met	Phe	Trp 300	Met	Ile	Gly	Gly	Val 305	Tyr	Ala	Ala	Ala	
_	-		-		atc Ile 315						_					1374
					cac His											1422
					gcc Ala			-	-		_	-	_		-	1470
	_				gag Glu			_		_		_			-	1518
_	_	_	_		gac Asp			_	_	-	_					1566
_	_			_	acc Thr 395				_			_		_	-	1614
_				_	tca Ser	_						_	_			1662
		-	_	-	ctg Leu			_	_							1710
-		_			tac Tyr		_			_	-	_	_		_	1758
					ttc Phe											1806
					ctc Leu 475											1854
					cgc Arg											1902
					cga Arg											1950
ggg	ctg	cgt	ctg	aag	tca	gtg	tcc	ttt	gag	gat	tcc	ctg	ttt	gag	gaa	1998

Gly	Leu	Arg 520	Leu	Lys	Ser	Val	Ser 525	Phe	Glu	Asp	Ser	Leu 530	Phe	Glu	Glu	
					gtc Val											2046
					gtg Val 555											2094
ttc Phe	gtg Val	aac Asn	agc Ser	cgc Arg 570	ctg Leu	gtg Val	aac Asn	agc Ser	aca Thr 575	ttc Phe	ctg Leu	cac His	aat Asn	aag Lys 580	gaa Glu	2142
					gtg Val											2190
					ttg Leu											2238
					atg Met											2286
					tcc Ser 635											2334
					atg Met											2382
					aac Asn											2430
cct Pro	tcc Ser	gac Asp 680	aag Lys	agg Arg	acg Thr	acg Thr	gcc Ala 685	ttc Phe	ggc Gly	ttc Phe	ctg Leu	aat Asn 690	gcc Ala	ctg Leu	tgt Cys	2478
aag Lys	ctg Leu 695	gca Ala	gct Ala	gta Val	ctg Leu	ggc Gly 700	atc Ile	agc Ser	atc Ile	ttc Phe	acg Thr 705	tcc Ser	ttt Phe	gtg Val	gga Gly	2526
atc Ile 710	acc Thr	aag Lys	gcc Ala	gct Ala	ccc Pro 715	atc Ile	ctc Leu	ttc Phe	gcc Ala	tca Ser 720	gct Ala	gcg Ala	ctt Leu	gcc Ala	ctt Leu 725	2574
					ctg Leu											2622
cag	tga	ggg	atgg	ggg (agtg	tctc	ag g	ggct	ttag	g ga	tggc	aggc	aca	ctgt	gac	2678

Gln

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cttagctgtg	tgcctgtgtg	catgtgtgtg	accctgacgg	gcaggggcta	cggggagggt	2798
ccctttgtcc	catgtttggg	aggagggact	ccccacctgc	tgccaccctc	aactttgcac	2858
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taccactata	ctgttcaatt	ataagccaag	agtagtagtt	tcagtgagca	cacacacaac	3338
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tactcttgtg	gctgctttct	ttggtactct	tcccactccc	accgtagctg	tgacgtgttg	3818
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<212> PRT

<213> Rattus norvegicus

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Asp Ile Ala Lys Glu Val Lys Lys His Ala Ala Lys Lys Val Val Lys

Gly Leu Asp Arg Val Gln Asp Glu Tyr Ser Arg Arg Ser Tyr Ser Arg Phe Glu Glu Glu Asp Asp Asp Phe Pro Ala Pro Ala Asp Gly Tyr Tyr Arg Gly Glu Gly Ala Gln Asp Glu Glu Gly Gly Ala Ser Ser Asp Ala Thr Glu Gly His Asp Glu Asp Asp Glu Ile Tyr Glu Gly Glu Tyr Gln Gly Ile Pro Arg Ala Glu Ser Gly Gly Lys Gly Glu Arg Met Ala Asp Gly Ala Pro Leu Ala Gly Val Arg Gly Gly Leu Ser Asp Gly Glu Gly Pro Pro Gly Gly Arg Gly Glu Ala Gln Arg Arg Lys Asp Arg Glu Glu Leu Ala Gln Gln Tyr Glu Thr Ile Leu Arg Glu Cys Gly His Gly Arg Phe Gln Trp Thr Leu Tyr Phe Val Leu Gly Leu Ala Leu Met Ala Asp Gly Val Glu Val Phe Val Val Gly Phe Val Leu Pro Ser Ala Glu Lys Asp Met Cys Leu Ser Asp Ser Asn Lys Gly Met Leu Gly Leu Ile Val Tyr Leu Gly Met Met Val Gly Ala Phe Leu Trp Gly Gly Leu Ala Asp Arg Leu Gly Arg Arg Gln Cys Leu Leu Ile Ser Leu Ser

Val Asn Ser Val Phe Ala Phe Phe Ser Ser Phe Val Gln Gly Tyr Gly

Thr Phe Leu Phe Cys Arg Leu Leu Ser Gly Val Gly Ile Gly Gly Ser Ile Pro Ile Val Phe Ser Tyr Phe Ser Glu Phe Leu Ala Gln Glu Lys Arg Gly Glu His Leu Ser Trp Leu Cys Met Phe Trp Met Ile Gly Gly Val Tyr Ala Ala Ala Met Ala Trp Ala Ile Ile Pro His Tyr Gly Trp Ser Phe Gln Met Gly Ser Ala Tyr Gln Phe His Ser Trp Arg Val Phe Val Leu Val Phe Ala Phe Pro Ser Val Phe Ala Ile Gly Ala Leu Thr Thr Gln Pro Glu Ser Pro Arg Phe Phe Leu Glu Asn Gly Lys His Asp Glu Ala Trp Met Val Leu Lys Gln Val His Asp Thr Asn Met Arg Ala Lys Gly His Pro Glu Arg Val Phe Ser Val Thr His Ile Lys Thr Ile His Gln Glu Asp Glu Leu Ile Glu Ile Gln Ser Asp Thr Gly Thr Trp Tyr Gln Arg Trp Gly Val Arg Ala Leu Ser Leu Gly Gly Gln Val Trp Gly Asn Phe Leu Ser Cys Phe Ser Pro Glu Tyr Arg Arg Ile Thr Leu Met Met Met Gly Val Trp Phe Thr Met Ser Phe Ser Tyr Tyr Gly Leu

Thr Val Trp Phe Pro Asp Met Ile Arg His Leu Gln Ala Val Asp Tyr

465					470					4/5					480	
Ala	Ala	Arg	Thr	Lys 485	Val	Phe	Pro	Gly	Glu 490	Arg	Val	Glu	His	Val 495	Thr	
Phe	Asn	Phe	Thr 500	Leu	Glu	Asn	Gln	Ile 505	His	Arg	Gly	Gly	Gln 510	Tyr	Phe	
Asn	Asp	Lys 515	Phe	Ile	Gly	Leu	Arg 520	Leu	Lys	Ser	Val	Ser 525	Phe	Glu	Asp	
Ser	Leu 530	Phe	Glu	Glu	Cys	Tyr 535	Phe	Glu	Asp	Val	Thr 540	Ser	Ser	Asn	Thr	
Phe 545	Phe	Arg	Asn	Cys	Thr 550	Phe	Ile	Asn	Thr	Val 555	Phe	Tyr	Asn	Thr	Asp 560	
Leu	Phe	Glu	Tyr	Lys 565	Phe	Val	Asn	Ser	Arg 570	Leu	Val	Asn	Ser	Thr 575	Phe	
Leu	His	Asn	Lys 580	Glu	Gly	Cys	Pro	Leu 585	Asp	Val	Thr	Gly	Thr 590	Gly	Glu	
Gly	Ala	Tyr 595	Met	Val	Tyr	Phe	Val 600	Ser	Phe	Leu	Gly	Thr 605	Leu	Ala	Val	
Leu	Pro 610	Gly	Asn	Ile	Val	Ser 615	Ala	Leu	Leu	Met	Asp 620	Lys	Ile	Gly	Arg	
Leu 625	Arg	Met	Leu	Ala	Gly 630	Ser	Ser	Val	Leu	Ser 635	Cys	Val	Ser	Cys	Phe 640	
Phe	Leu	Ser	Phe	Gly 645	Asn	Ser	Glu	Ser	Ala 650	Met	Ile	Ala	Leu	Leu 655	Cys	
Leu	Phe	Gly	Gly 660	Val	Ser	Ile	Ala	Ser 665	Trp	Asn	Ala	Leu	Asp 670	Val	Leu	
Thr	Val	Glu 675	Leu	Tyr	Pro	Ser	Asp 680	Lys	Arg	Thr	Thr	Ala 685	Phe	Gly	Phe	
Len	Aen	Δ 1 =	T All	Cue	Tare	T.O.II	Δla	Δla	Val	T.011	Clv	Tla	Ser	Tle	Phe	

690 695 700 Thr Ser Phe Val Gly Ile Thr Lys Ala Ala Pro Ile Leu Phe Ala Ser 715 705 710 Ala Ala Leu Ala Leu Gly Ser Ser Leu Ala Leu Lys Leu Pro Glu Thr 725 730 Arg Gly Gln Val Leu Gln 740 <210> 11 <211> 2052 <212> DNA <213> Rattus norvegicus <220> <221> CDS <222> (1)..(2052)<400> 11 atg gat gac tac agg tat cgg gac aac tat gag ggc tat gcc cct aat Met Asp Asp Tyr Arg Tyr Arg Asp Asn Tyr Glu Gly Tyr Ala Pro Asn gat ggc tac tac cgg ggc aat gag cag aac ccg gaa gaa gat gca cag 96 Asp Gly Tyr Tyr Arg Gly Asn Glu Gln Asn Pro Glu Glu Asp Ala Gln 20 144 age gat gtt aca gaa gge cae gat gaa gag gat gag ate tat gag gge Ser Asp Val Thr Glu Gly His Asp Glu Glu Asp Glu Ile Tyr Glu Gly 40 35 gag tac caa ggc atc cct cat cca gat gat gtc aag tct aag cag act 192 Glu Tyr Gln Gly Ile Pro His Pro Asp Asp Val Lys Ser Lys Gln Thr 55 aag atg gca ccg tcc aga gca gat ggc ctt cgg ggc cag gca gac ctg 240 Lys Met Ala Pro Ser Arg Ala Asp Gly Leu Arg Gly Gln Ala Asp Leu 70 atg gct gag aga atg gaa gat gag gag cag ctc gct cac cag tac gag 288 Met Ala Glu Arg Met Glu Asp Glu Glu Gln Leu Ala His Gln Tyr Glu 85 90 336 acc atc att gat gag tgt ggc cat ggg cgc ttc cag tgg acc ctc ttt Thr Ile Ile Asp Glu Cys Gly His Gly Arg Phe Gln Trp Thr Leu Phe 100 384 ttc gtc ttg gtc ttg gcc ttg atg gct gac gga gtg gaa gtg ttt gtg

Phe Val Leu Val Leu Ala Leu Met Ala Asp Gly Val Glu Val Phe Val

115		120	125		
	ct ctg cca agt la Leu Pro Ser 135			, ,	32
= -	ga atg ctc ggg ly Met Leu Gly 150			, , J	80
	tc ctg ggg ggc le Leu Gly Gly 165			, , ,	28
Val Leu Ser Me	tg tcc ttg gct et Ser Leu Ala 80			Leu Ser	76
	ag gga tat gga ln Gly Tyr Gly				24
	tt ggg ggc tcc le Gly Gly Ser 215				72
	ca cgg gag aaa er Arg Glu Lys 230	Arg Gly Glu		, , , , ,	20
	tg act ggg ggc et Thr Gly Gly 245				68
Ile Ile Pro H	ac tat ggc tgg is Tyr Gly Trp 60			n Tyr His	16
	gg aga gtg ttt rp Arg Val Phe				64
	tg gcc ctg aag al Ala Leu Lys 295	_		,	12
	gc aag cat gat ly Lys His Asp 310	Glu Ala Trp	_	,	60
	ac atg aga gct sn Met Arg Ala 325				80
	tc aaa act ccc le Lys Thr Pro				56

340	345	350
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	aac gcc ttg tac t Asn Ala Leu Tyr C 380	
	ctg gct gtg gtc t Leu Ala Val Val I 395	
Ala Leu Ser Tyr	gtg tgg ttc ccc g Val Trp Phe Pro F 410	
	tct aaa atg aag g Ser Lys Met Lys V 425	
	aac ttc acc atg c Asn Phe Thr Met G	
	gat aag ttc ata a Asp Lys Phe Ile I 460	
	ttc ttt gac aaa t Phe Phe Asp Lys C 475	
Asp Val Thr Ser	ttc aag aac tgc a Phe Lys Asn Cys T 490	
	tac aaa cac aag t Tyr Lys His Lys F 505	
	gag cag aag gag g Glu Gln Lys Glu G	
	ctg att tac ctc of Leu Ile Tyr Leu V 540	
	aac ata att tct c Asn Ile Ile Ser A 555	
	att ggt ggc tcc a Ile Gly Gly Ser N	

565		570	575
gca gtc tgc tgc ttc Ala Val Cys Cys Phe 580		Gly Asn Ser Glu S	
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gct ctg gat gtg atc Ala Leu Asp Val Ile 610			
act gcc ttc ggc atc Thr Ala Phe Gly Ile 625			
gga aac act atc ttt Gly Asn Thr Ile Phe 645			
atc ctt ctg gct gct Ile Leu Leu Ala Ala 660		Gly Gly Gly Leu V	-
cga ctg cca gag act Arg Leu Pro Glu Thr 675			2052
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Ser Asp Val Thr Glu			
35	Gly His Asp Glu 40	Glu Asp Glu Ile T 45	yr Glu Gly
	40	45	·

Met Ala Glu Arg Met Glu Asp Glu Glu Gln Leu Ala His Gln Tyr Glu 85 90 95

Thr Ile Ile Asp Glu Cys Gly His Gly Arg Phe Gln Trp Thr Leu Phe 100 105 110

Phe Val Leu Val Leu Ala Leu Met Ala Asp Gly Val Glu Val Phe Val 115 120 125

Val Ser Phe Ala Leu Pro Ser Ala Glu Lys Asp Met Cys Leu Ser Ser 130 135 140

Ser Lys Lys Gly Met Leu Gly Leu Ile Val Tyr Leu Gly Met Met Ala 145 150 155 160

Gly Ala Phe Ile Leu Gly Gly Leu Ala Asp Lys Leu Gly Arg Lys Lys 165 170 175

Val Leu Ser Met Ser Leu Ala Ile Asn Ala Ser Phe Ala Ser Leu Ser 180 185 190

Ser Phe Val Gln Gly Tyr Gly Ala Phe Leu Phe Cys Arg Leu Ile Ser 195 200 205

Gly Ile Gly Ile Gly Gly Ser Leu Pro Ile Val Phe Ala Tyr Phe Ser 210 215 220

Glu Phe Leu Ser Arg Glu Lys Arg Gly Glu His Leu Ser Trp Leu Gly 225 230 235 240

Ile Phe Trp Met Thr Gly Gly Ile Tyr Ala Ser Ala Met Ala Trp Ser 245 250 255

Ile Ile Pro His Tyr Gly Trp Gly Phe Ser Met Gly Thr Asn Tyr His 260 265 270

Phe His Ser Trp Arg Val Phe Val Ile Val Cys Ala Leu Pro Ala Thr 275 280 285

Val Ser Met Val Ala Leu Lys Phe Met Pro Glu Ser Pro Arg Phe Leu 290 295 300

Leu 305	Glu	Met	Gly	Lys	His 310	Asp	Glu	Ala	Trp	Met 315	Ile	Leu	Lys	Gln	Val 320
His	Asp	Thr	Asn	Met 325	Arg	Ala	Lys	Gly	Thr 330	Pro	Glu	Lys	Val	Phe 335	Thr
Val	Ser	His	Ile 340	Lys	Thr	Pro	Lys	Gln 345	Met	Asp	Glu	Phe	Ile 350	Glu	Ile
Gln	Ser	Ser 355	Thr	Gly	Thr	Trp	Tyr 360	Gln	Arg	Trp	Leu	Val 365	Arg	Phe	Met
Thr	Ile 370	Phe	Lys	Gln	Val	Trp 375	Asp	Asn	Ala	Leu	Tyr 380	Cys	Val	Met	Gly
Pro 385	Tyr	Arg	Met	Asn	Thr 390	Leu	Ile	Leu	Ala	Val 395	Val	Trp	Phe	Thr	Met 400
Ala	Leu	Ser	Tyr	Tyr 405	Gly	Leu	Thr	Val	Trp 410	Phe	Pro	Asp	Met	Ile 415	Arg
Tyr	Phe	Gln	Asp 420	Glu	Glu	Tyr	Lys	Ser 425	Lys	Met	Lys	Val	Phe 430	Phe	Gly
Glu	His	Val 435	His	Gly	Ala	Thr	Ile 440	Asn	Phe	Thr	Met	Glu 445	Asn	Gln	Ile
His	Gln 450	His	Gly	Lys	Leu	Val 455	Asn	Asp	Lys	Phe	Ile 460	Lys	Met	Tyr	Phe
Lys 465	His	Val	Leu	Phe	Glu 470	Asp	Thr	Phe	Phe	Asp 475	Lys	Cys	Tyr	Phe	Glu 480
Asp	Val	Thr	Ser	Thr 485	Asp	Thr	Tyr	Phe	Lys 490	Asn	Cys	Thr	Ile	Glu 495	Ser
Thr	Thr	Phe	Tyr 500	Asn	Thr	Asp	Leu	Tyr 505	Lys	His	Lys	Phe	Ile 510	Asp	Cys
Arg	Phe	Ile 515	Asn	Ser	Thr	Phe	Leu 520	Glu	Gln	Lys	Glu	Gly 525	Cys	His	Met

Asp Phe Glu Glu Asp Asn Asp Phe Leu Ile Tyr Leu Val Ser Phe Leu 530 Gly Ser Leu Ser Val Leu Pro Gly Asn Ile Ile Ser Ala Leu Leu Met 550 545 Asp Arg Ile Gly Arg Leu Lys Met Ile Gly Gly Ser Met Leu Ile Ser 565 570 Ala Val Cys Cys Phe Phe Leu Phe Phe Gly Asn Ser Glu Ser Ala Met 580 585 Ile Gly Trp Gln Cys Leu Phe Cys Gly Thr Ser Ile Ala Ala Trp Asn 595 600 Ala Leu Asp Val Ile Thr Val Glu Leu Tyr Pro Thr Asn Gln Arg Ala 610 615 Thr Ala Phe Gly Ile Leu Asn Gly Leu Cys Lys Leu Gly Ala Ile Leu 635 Gly Asn Thr Ile Phe Ala Ser Phe Val Gly Ile Thr Lys Val Val Pro 645 650 Ile Leu Leu Ala Ala Ser Leu Val Gly Gly Leu Val Ala Leu 660 665 Arg Leu Pro Glu Thr Arg Glu Gln Val Leu Met 675 <210> 13 <211> 2622 <212> DNA <213> Rattus norvegicus <220> <221> CDS <222> (224)..(2407) <400> 13 gcgcgctgca ggaagagtgg cagaccgaag cggcctcggg ctgcaaacgg aggggcgctc 60 120 qcqcqqcqac qqctqcaqqq ctqacaccqc tcagggcagg ggggtcccag gcggctggaa

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gtggactctt c	cctgctgtg c	cttgcccat cg	gccgagat		gac tcc Asp Ser	235
tac aag gat Tyr Lys Asp 5						283
gag gtg aag Glu Val Lys						331
gcc cag gat Ala Gln Asp	-		_	-		379
gat gat gat Asp Asp Asp 55						427
gcc aat gat Ala Asn Asp 70			-			475
gaa gac gag Glu Asp Glu 85						523
caa ggg aag Gln Gly Lys						571
tac aag gac Tyr Lys Asp						619
gag ctc gcc Glu Leu Ala 135						667
cgt ttc cag Arg Phe Gln 150		_				715
gac ggc gtg Asp Gly Val 165				_		763
aca gac cta Thr Asp Leu						811
gtg tac ctc	ggg atg atg	gtg ggg gcg	ttc ttc	tgg gga gga	ctg gca	859

Val	Tyr	Leu	Gly 200	Met	Met	Val	Gly	Ala 205	Phe	Phe	Trp	Gly	Gly 210	Leu	Ala	
					aag Lys											907
					ctt Leu											955
					ctt Leu 250											1003
					ttt Phe											1051
					ctc Leu											1099
					tgg Trp											1147
					tac Tyr											1195
_	_	-		_	tgc Cys 330	_		_			-					1243
					ttc Phe											1291
	_		_	_	cta Leu			-			-	-	-			1339
					ttc Phe											1387
					gag Glu											1435
					atc Ile 410											1483
ttt	atg	aga	tgc	ttc	aac	tac	ccg	gtc	agg	gaa	aac	acc	ata	aag	ctt	1531

Phe	Met	Arg	Cys	Phe 425	Asn	Tyr	Pro	Val	Arg 430	Glu	Asn	Thr	Ile	Lys 435	Leu		
										tac Tyr						1579	
										tct Ser						1627	
										gca Ala						1675	
										atg Met 495						1723	
										acc Thr						1771	
	_		_			_	_			tca Ser	-					1819	
										gag Glu						1867	
ccc Pro	tat Tyr 550	aaa Lys	ttc Phe	ata Ile	gac Asp	agc Ser 555	gag Glu	ttt Phe	caa Gln	aac Asn	tgc Cys 560	tcg Ser	ttt Phe	ctt Leu	cac His	1915	
										gac Asp 575						1963	
										ttg Leu						2011	
										atc Ile						2059	
cta Leu	ggt Gly	ggc Gly 615	tcc Ser	atg Met	gtg Val	ctc Leu	tcg Ser 620	ggg Gly	atc İle	agc Ser	tgc Cys	ttc Phe 625	ttc Phe	ctg Leu	tgg Trp	2107	
										atg Met						2155	
gga	ctg	acc	atc	tca	gcg	tgg	aac	tct	ctt	gat	gta	gtc	acg	gtg	gaa	2203	

Gly Leu Thr Ile Ser Ala Trp Asn Ser Leu Asp Val Val Thr Val Glu 645 650 655 660	
cta tac ccc aca gac cgg aga gca acg ggc ttt ggc ttc ttg aac gca Leu Tyr Pro Thr Asp Arg Arg Ala Thr Gly Phe Gly Phe Leu Asn Ala 665 670 675	2251
ctc tgt aaa gca gcg gcc gtc ctg gga aac tta ata ttc ggc tcc ttg Leu Cys Lys Ala Ala Ala Val Leu Gly Asn Leu Ile Phe Gly Ser Leu 680 685 690	2299
gtc agc atc acc aaa gca atc cct atc ctg ctg gct tcc acc gtg ctc Val Ser Ile Thr Lys Ala Ile Pro Ile Leu Leu Ala Ser Thr Val Leu 695 700 705	2347
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gtt ctg atg tga caaaagccat tctcttctct caccatgggt cagccctatt Val Leu Met 725	2447
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Asp Ile Ala Lys Glu Val Lys Lys Gln Thr Val Lys Lys Val Asn Gln 20 25 30 Ala Val Asp Arg Ala Gln Asp Glu Tyr Thr Gln Arg Ser Tyr Ser Arg	

Gly His	Asp	Glu	Glu	Asp	Glu	Ile	Tyr	Glu	Gly	Glu	Tyr	Gln	Gly	Ile
			85					90					95	

Pro Ser Thr Asn Gln Gly Lys Asp Ser Ile Val Ser Val Gly Gln Pro 100 105 110

Lys Gly Asp Glu Tyr Lys Asp Arg Arg Glu Leu Glu Ser Glu Arg Arg 115 120 125

Ala Asp Glu Glu Glu Leu Ala Gln Gln Tyr Glu Leu Ile Ile Gln Glu 130 135 140

Cys Gly His Gly Arg Phe Gln Trp Ala Leu Phe Phe Val Leu Gly Met 145 150 155 160

Ala Leu Met Ala Asp Gly Val Glu Val Phe Val Val Gly Phe Val Leu 165 170 175

Pro Ser Ala Glu Thr Asp Leu Cys Ile Pro Asn Ser Gly Ser Gly Trp 180 185 190

Leu Gly Ser Ile Val Tyr Leu Gly Met Met Val Gly Ala Phe Phe Trp 195 200 205

Gly Gly Leu Ala Asp Lys Val Gly Arg Lys Gln Ser Leu Leu Ile Cys 210 215 220

Met Ser Val Asn Gly Phe Phe Ala Phe Leu Ser Ser Phe Val Gln Gly 225 230 235 240

Tyr Gly Phe Phe Leu Leu Cys Arg Leu Leu Ser Gly Phe Gly Ile Gly 245 250 255

Gly Ala Ile Pro Thr Val Phe Ser Tyr Phe Ala Glu Val Leu Ala Arg 260 265 270

Glu Lys Arg Gly Glu His Leu Ser Trp Leu Cys Met Phe Trp Met Ile 275 280 285

Gly Gly Ile Tyr Ala Ser Ala Met Ala Trp Ala Ile Ile Pro His Tyr 290 295 300

Gly 305	Trp	Ser	Phe	Ser	Met 310	Gly	Ser	Ala	Tyr	Gln 315	Phe	His	Ser	Trp	Arg 320
Val	Phe	Val	Ile	Val 325	Cys	Ala	Leu	Pro	Cys 330	Val	Ser	Ser	Val	Val 335	Ala
Leu	Thr	Phe	Met 340	Pro	Glu	Ser	Pro	Arg 345	Phe	Leu	Leu	Glu	Val 350	Gly	Lys
His	Asp	Glu 355	Ala	Trp	Met	Ile	Leu 360	Lys	Leu	Ile	His	Asp 365	Thr	Asn	Met
Arg	Ala 370	Arg	Gly	Gln	Pro	Glu 375	Lys	Val	Phe	Thr	Val 380	Asn	Lys	Ile	Lys
Thr 385	Pro	Lys	Gln	Ile	Asp 390	Glu	Leu	Ile	Glu	Ile 395	Glu	Ser	Asp	Thr	Gly 400
Thr	Trp	Tyr	Arg	Arg 405	Cys	Phe	Val	Arg	Ile 410	Arg	Thr	Glu	Leu	Tyr 415	Gly
Ile	Trp	Leu	Thr 420	Phe	Met	Arg	Cys	Phe 425	Asn	Tyr	Pro	Val	Arg 430	Glu	Asn
Thr	Ile	Lys 435	Leu	Thr	Ile	Val	Trp 440	Phe	Thr	Leu	Ser	Pḥe 445	Gly	Tyr	Tyr
Gly	Leu 450	Ser	Val	Trp	Phe	Pro 455	Asp	Val	Ile	Lys	His 460	Leu	Gln	Ser	Asp
Glu 465	Tyr	Ala	Leu	Leu	Thr 470	Arg	Asn	Val	Gln	Lys 475	Asp	Lys	Tyr	Ala	Asn 480
Phe	Ser	Ile	Asn	Phe 485	Thr	Met	Glu	Asn	Gln 490	Val	His	Thr	Gly	Met 495	Glu
Tyr	Asp	Asn	Gly 500	Arg	Phe	Leu	Gly	Val 505	Lys	Phe	Lys	Ser	Val 510	Thr	Phe
Lys	Asp	Ser 515	Val	Phe	Lys	Ser	Cys 520	Thr	Phe	Asp	Asp	Val 525	Thr	Ser	Val

Asn Thr Tyr Phe Lys Asn Cys Thr Phe Ile Asp Thr Leu Phe Glu Asn 530 535 540

Thr Asp Phe Glu Pro Tyr Lys Phe Ile Asp Ser Glu Phe Gln Asn Cys 545 550 555 560

Ser Phe Leu His Asn Lys Thr Gly Cys Gln Ile Thr Phe Asp Asp Asp 565 570 575

Tyr Ser Ala Tyr Trp Ile Tyr Phe Val Asn Phe Leu Gly Thr Leu Ala 580 585 590

Val Leu Pro Gly Asn Ile Val Ser Ala Leu Leu Met Asp Arg Ile Gly 595 600 605

Arg Leu Thr Met Leu Gly Gly Ser Met Val Leu Ser Gly Ile Ser Cys 610 615 620

Phe Phe Leu Trp Phe Gly Thr Ser Glu Ser Met Met Ile Gly Met Leu 625 630 635 640

Cys Leu Tyr Asn Gly Leu Thr Ile Ser Ala Trp Asn Ser Leu Asp Val 645 650 655

Val Thr Val Glu Leu Tyr Pro Thr Asp Arg Arg Ala Thr Gly Phe Gly 660 665 670

Phe Leu Asn Ala Leu Cys Lys Ala Ala Ala Val Leu Gly Asn Leu Ile 675 680 685

Phe Gly Ser Leu Val Ser Ile Thr Lys Ala Ile Pro Ile Leu Leu Ala 690 695 700

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						gct Ala 55								19	92
						gaa Glu								24	40
_		_		_		tct Ser	-			_	-			28	38
_	-	_		_	_	att Ile	_	-	_			_	_	33	36
_			_			agc Ser								38	34
						tcc Ser 135								43	32
						aca Thr								48	80
						agc Ser								52	28
						gtg Val								57	76

		tac Tyr											624
		att Ile											672
		gtg Val											720
		gcc Ala 245											768
		agt Ser											816
		acc Thr											864
		aag Lys											912
		ttc Phe											960
		ttc Phe 325											1008
	Glu	ctc Leu	Phe	Gln	Ala	Gly	Asp	Val	Cys	Ser	Ile	Ser	1056
		gtg Val											1104
		tac Tyr											1152
		gtg Val											1200
		ctg Leu 405											1248

ctg ttc atc tgc at Leu Phe Ile Cys Il 420			
gca aga gcg ttt at Ala Arg Ala Phe Il 435		Phe Gln Ala Ala '	
cct gag gtg tat cc Pro Glu Val Tyr Pr 450			
agc ggc atg gcg ag Ser Gly Met Ala Ar 465			
gtg atg ctg gaa tc Val Met Leu Glu Se 48	r Ser Val Tyr		
tgc tgc ctc ctt gc Cys Cys Leu Leu Al 500			
aaa ggc cga gca ct Lys Gly Arg Ala Le 515		Ser His Arg Glu '	
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Arg Arg Thr Gly Gl 20	u Ser Ala Arg	Ser Glu Asp Asp 2 25	Ala Ala Ser Gly 30
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Glu Leu Asp Asp Gly Ala Ala Val Pro Lys Glu Phe Ala Asn Pro Thr 50 55 60

Asp Asp Thr Phe Met Val Glu Asp Ala Val Glu Ala Ile Gly Phe Gly 65 70 75 80

Arg Phe Gln Trp Lys Leu Ser Val Leu Thr Gly Leu Ala Trp Met Ala 85 90 95

Asp Ala Met Glu Met Met Ile Leu Ser Ile Leu Ala Pro Gln Leu His 100 105 110

Cys Glu Trp Arg Leu Pro Ser Trp Gln Val Ala Leu Leu Thr Ser Val 115 120 125

Val Phe Ile Gly Met Met Ser Ser Ser Thr Leu Trp Gly Asn Ile Ser 130 135 140

Asp Gln Tyr Gly Arg Lys Thr Gly Leu Lys Ile Ser Val Phe Trp Thr 145 150 155 160

Leu Tyr Tyr Gly Ile Leu Ser Ala Phe Ala Pro Val Tyr Ser Trp Ile 165 170 175

Leu Val Leu Arg Gly Leu Val Gly Phe Gly Ile Gly Gly Val Pro Gln
180 185 190

Ser Val Thr Leu Tyr Ala Glu Phe Leu Pro Met Lys Ala Arg Ala Lys 195 200 205

Cys Ile Leu Leu Ile Glu Val Phe Trp Ala Ile Gly Thr Val Phe Glu 210 215 220

Val Leu Leu Ala Val Phe Val Met Pro Ser Leu Gly Trp Arg Trp Leu 225 230 235 240

Leu Leu Ser Ala Ala Pro Leu Leu Val Phe Ala Val Leu Cys Phe 245 250 255

Trp Leu Pro Glu Ser Ala Arg Tyr Asp Val Leu Ser Gly Asn Gln Glu 260 265 270

Lys Ala Ile Ala Thr Leu Lys Arg Ile Ala Thr Glu Asn Gly Ala Pro $275 \hspace{1.5cm} 280 \hspace{1.5cm} 285$

Met Pro Leu Gly Lys Leu Ile Ile Ser Arg Gln Glu Asp Arg Gly Lys 290 295 300

Met Arg Asp Leu Phe Thr Pro His Phe Arg Trp Thr Thr Leu Leu Leu 305 310 315 320

Trp Phe Ile Trp Phe Ser Asn Ala Phe Ser Tyr Tyr Gly Leu Val Leu 325 330 335

Leu Thr Thr Glu Leu Phe Gln Ala Gly Asp Val Cys Ser Ile Ser Ser 340 345 350

Arg Lys Lys Ala Val Glu Ala Lys Cys Ser Leu Ala Cys Glu Tyr Leu 355 360 . 365

Ser Lys Glu Asp Tyr Met Asp Leu Leu Trp Thr Thr Leu Ser Glu Phe 370 375 380

Pro Gly Val Leu Val Thr Leu Trp Val Ile Asp Arg Leu Gly Arg Lys 385 390 395 400

Lys Thr Met Ala Leu Cys Phe Val Ile Phe Ser Leu Cys Ser Leu Leu 405 410 415

Leu Phe Ile Cys Ile Gly Arg Asn Val Leu Thr Leu Leu Phe Ile 420 425 430

Ala Arg Ala Phe Ile Ser Gly Gly Phe Gln Ala Ala Tyr Val Tyr Thr $435 \hspace{1.5cm} 440 \hspace{1.5cm} 445 \hspace{1.5cm}$

Pro Glu Val Tyr Pro Thr Ala Thr Arg Ala Leu Gly Leu Gly Thr Cys 450 455 460

Ser Gly Met Ala Arg Val Gly Ala Leu Ile Thr Pro Phe Ile Ala Gln 465 470 475 480

Val Met Leu Glu Ser Ser Val Tyr Leu Thr Leu Ala Val Tyr Ser Gly 485 490 495

Cys Cys Leu Leu Ala Ala Leu Ala Ser Cys Phe Leu Pro Ile Glu Thr 500 505 510

Lys Gly Arg Ala Leu Gln Glu Ser Ser His Arg Glu Trp Gly Gln Glu 515 520 525

Met Val Gly Arg Gly Thr Asn Ser Thr Gly Val Pro Arg Ser Asn Ser 530 535 540

Gly Ser Gln Glu 545